'PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION *US/08/436,339*

1806

DATE: 10/15/97
TIME: 12:39:25

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> This Raw Listing contains the General
> Information Section and up to the first 5 pages.

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 1                              SEQUENCE LISTING
 2
 3   (1)     General Information:
 4
 5      (i) APPLICANT: Papayannopoulou, Thalia
 6
 7     (ii) TITLE OF INVENTION:  Peripheralization of Hematopoietic
 8                               Stem Cells
 9
10    (iii) NUMBER OF SEQUENCES: 2
11
12     (iv) CORRESPONDENCE ADDRESS:
13          (A) ADDRESSEE: Allegretti & Witcoff, Ltd.
14          (B) STREET: 75 State Street
15          (C) CITY: Boston
16          (D) STATE: Massachusetts
17          (E) COUNTRY: USA
18          (F) ZIP: 02109
19
20      (v) COMPUTER READABLE FORM:
21          (A) MEDIUM TYPE: Floppy disk
22          (B) COMPUTER: IBM PC compatible
23          (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24          (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
25
26     (vi) CURRENT APPLICATION DATA:
27          (A) APPLICATION NUMBER:  08/436,339
28          (B) FILING DATE: 13-JUL-1995
29          (C) CLASSIFICATION: 424
30
31    (vii) PRIOR APPLICATION DATA:
32          (A) APPLICATION NUMBER: 07/977,702
33          (B) FILING DATE: 13-NOV-1992
34
35   (viii) ATTORNEY/AGENT INFORMATION:
36          (A) NAME: Keown, Wayne A
37          (B) REGISTRATION NUMBER: 33,923
38          (C) REFERENCE/DOCKET NUMBER: 92,678; D014
39
40     (ix) TELECOMMUNICATION INFORMATION:
41          (A) TELEPHONE: 617/345-9100
42          (B) TELEFAX: 617/345-9111
43
44
45   (2) INFORMATION FOR SEQ ID NO:1:
46
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47          (i) SEQUENCE CHARACTERISTICS:
48              (A) LENGTH: 360 base pairs
49              (B) TYPE: nucleic acid
50              (C) STRANDEDNESS: single
51              (D) TOPOLOGY: linear
52
53      (ii) MOLECULE TYPE: cDNA
54
55     (iii) HYPOTHETICAL: NO
56
57      (iv) ANTI-SENSE: NO
58
59      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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62  GTCAAACTGC AGCAGTCTGG GGCAGAGCTT GTGAAGCCAG GGGCCTCAGT CAAGTTGTCC    60
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64  TGCACAGCTT CTGGCTTCAA CATTAAAGAC ACCTATATGC ACTGGGTGAA GCAGAGGCCT   120
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70  CAGCTCAGCA GCCTGACATC TGAGGACACT GCCGTCTACT ACTGTGCAGA CGGAATGTGG   300
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72  GTATCAACGG GATATGCTCT GGACTTCTGG GGCCAAGGGA CCACGGTCAC CGTCTCCTCA   360
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77  (2) INFORMATION FOR SEQ ID NO:2:
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79          (i) SEQUENCE CHARACTERISTICS:
80              (A) LENGTH: 318 base pairs
81              (B) TYPE: nucleic acid
82              (C) STRANDEDNESS: single
83              (D) TOPOLOGY: linear
84
85      (ii) MOLECULE TYPE: cDNA
86
87     (iii) HYPOTHETICAL: NO
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89      (iv) ANTI-SENSE: NO
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91      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
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98  GGGCAGTCTC CTAAACTGCT GATATATTAT GCATCCAATC GCTACACTGG AGTCCCTGAT   180
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RAW SEQUENCE LISTING
PATENT APPLICATION *US/08/436,339*

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104   GGGACCAAGC TGGAGATC                                                    318
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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION   *US/08/436,339*

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